|    | CRF crors Corrected by the STIC Systems Branch  |
|----|---|
| J. | umber: 09/904,956 ENTED Edited by: 1  |
|    | changed a file from non-ASCII to ASCII  CRF Processing Dat : 2/6/200  CRF Processing Dat : 2/6/200  Control of the control of |
|    | Changed the margins in cases where the sequence text was "wrapped" down to the next line.   |
|    | Edited a format error in the Current Application Data section, specifically:  |
|    | Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other   |
|    | Added the mandatory heading and subheadings for "Current Application Data".   |
|    | Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.   |
| ,  | Changed the spelling of a mandatory field (the headings or subheadings), specifically:  |
| •  | Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:   |
|    | Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: /73  |
|    | Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  |
|    | Inserted colons after headings/subheadings. Headings edited included:   |
|    | Deleted extra, invalid, headings used by an applicant, specifically:  |
|    | Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of fi page numbers throughout text; other invalid text, such as  |
|    | Inserted mandatory headings, specifically:  |
|    | Corrected an obvious error in the response, specifically:   |
| •  | Edited identifiers where upper case is used but lower case is required, or vice versa.  |
|    | Corrected an error in the Number of Sequences field, specifically:  |
| -  | A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  |
|    | Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error lue to a Patentin bug). Sequences corrected:   |
|    | Other:  |
| -  | · · · · · · · · · · · · · · · · · · ·   |
|    | The above corrections must be communicated to the applicant in the first Offic  |

Examiner: The above corrections must be communicated to the applicant in the first Offic Action. DO NOT send a copy of this form.

3/1/95

#Y



OIPE

RAW SEQUENCE LISTING

3 <110> APPLICANT: Genentech, Inc.

DATE: 02/06/2002

PATENT APPLICATION: US/09/904,956

TIME: 08:16:11

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Output Set: N:\CRF3\02062002\I904956.raw

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              Eaton, Dan L.
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              Ferrara, Napoleone
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              Fong, Sherman
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              Hillan, Kenneth, J.
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              Kljavin, Ivar J.
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              Mather, Jennie P.
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              Pan, James
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     29 <120> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
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     32 <130> FILE REFERENCE: 10466-14
C--> 34 <140> CURRENT APPLICATION NUMBER: US/09/904,956
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     38 <151> PRIOR FILING DATE: 2000-02-22
     40 <150> PRIOR APPLICATION NUMBER: US 60/143,048
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DATE: 02/06/2002

TIME: 08:16:11

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|     |   |       |          |        |            | 7 200    | ת 1 ת      | 7 1 n      | T 011 | C1,,  | T 011        | T 011                | Dro        | T 011       | T 011    | Tou       |      |
|     |   | AIG   | Leu      | PIO    | AIG        | AIG      | Ата        | Ата        | ьец   |       | ьeu          | ьeu                  | PIO        | ьец         | Leu      | ьец       |      |
| 135 | 1   | T 0   | D        | D == 0 | כוג        | Dmo      | c1         | 7 ] n      | 7 l n | 10    | T            | Dwo                  | mh m       | Dwo         | 15       | TI i a    |      |
|     | ьeu   | ьeu   | Pro      |        | Ата        | PIO      | GLU        | Ald        |       | гуѕ   | гуѕ          | PIO                  | THE        |             | Cys      | HIS       |      |
| 138 |   | G     | 3        | 20     | <b>v</b>   | ¥7- 7    |            | <b>.</b>   | 25    |       | <b>G</b> 3   | <b>01</b>            | W-+        | 30          | 3        | m l       |      |
|     | Arg   | Cys   | _        | GLY    | ьeu        | vaı      | Asp        | _          | Pne   | ASN   | GIn          | GIY                  |            | val         | Asp      | Thr       |      |
| 141 | . 1 .   | T     | 35       |        | nl         | G 1      | <b>a</b> 1 | 40         | •     | m 1   | <b>3.</b> 1. | m                    | 45         | <b>a</b> 1  | <b>T</b> | m l       |      |
|     | Ата   |       | Lys      | Asn    | Pne        | GTÄ      |            | GTÄ        | Asn   | Tnr   | Ата          |                      | GIU        | GIU         | Lys      | Thr       |      |
| 144 | <b>.</b>  | 50    | <b>.</b> | _      | <b>a</b> 1 | <b>a</b> | 55         | <b>a</b> 1 | ~1.   |       | <b>.</b> .   | 60                   | <b>a</b> 1 | <b>-1</b> - | Ŧ        | <b>01</b> |      |
|     |   | Ser   | ьуs      | Tyr    | GIU        |          | ser        | GLu        | тте   | Arg   |              | Leu                  | GIU        | ше          | Leu      |           |      |
| 147 | 65  | _     | _        | -1     | _          | 70       | _          | -1         | ~1    | _     | 75           | ~ 1                  |            | _           | ~ 1      | 80        |      |
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| 150 | ~ 7   |       |          |        | 85         | - 3      |            |            | _     | 90    |              | _                    | _          | _           | 95       | _         |      |
|     | GIn   | GIu   | GLu      |        | Leu        | Glu      | Ala        | Trp        | _     | Leu   | GIn          | Leu                  | ьys        |             | Glu      | Tyr       |      |
| 153 | _   | _     | _        | 100    |            | _        | _,         | _          | 105   |       | _,           | _                    | _          | 110         | _        | _         |      |
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| 162 |   |       |          | _      |            | 150      |            |            |       |       | 155          |                      | _          | _           |          | 160       |      |
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| 168 |   |       |          | 180    |            |          |            |            | 185   |       |              |                      |            | 190         |          |           |      |
|     | His   | Ser   |          | Cys    | Thr        | Ala      | Cys        |            | Glu   | Ser   | Cys          | Lys                  | Thr        | Cys         | Ser      | Gly       |      |
| 171 |   |       | 195      |        |            |          |            | 200        |       |       |              |                      | 205        |             |          |           |      |
| 173 | Leu   |       | Asn      | Arg    | Asp        | Cys      | Gly        | Glu        | Cys   | Glu   | Val          | Gly                  | Trp        | Val         | Leu      | Asp       |      |
| 174 |   | 210   |          |        |            |          | 215        |            |       |       |              | 220                  |            |             |          |           |      |
| 176 | Glu   | Gly   | Ala      | Cys    | Val        | Asp      | Val        | Asp        | Glu   | Cys   | Ala          | Ala                  | Glu        | Pro         | Pro      | Pro       |      |
| 177 | 225   |       |          |        |            | 230      |            |            |       |       | 235          |                      |            |             |          | 240       |      |
|     | Cys   | Ser   | Ala      | Ala    | Gln        | Phe      | Cys        | Lys        | Asn   | Ala   | Asn          | Gly                  | Ser        | Tyr         | Thr      | Cys       |      |
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| 182 | Glu   | Glu   | Cys      | Asp    | Ser        | Ser      | Cys        | Val        | Gly   | Cys   | Thr          | Gly                  | Glu        | Gly         | Pro      | Gly       |      |
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| 185 | Asn   | Cys   | Lys      | Glu    | Cys        | Ile      | Ser        | Gly        | Tyr   | Ala   | Arg          | $\operatorname{Glu}$ | His        | Gly         | Gln      | Cys       |      |
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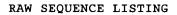


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| 189 | 290                            |           | 295       |         | _          |      | 300       | _     |       | _     | _      |      |  |  |
|     | Asn Glu As                     | n Cys Tyr |           | Pro GI  | ser        |      | Val       | Cys   | val   | Cys   |        |      |  |  |
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| 195 |                                | 325       |           | _       | 330        |      |           |       |       | 335   |        |      |  |  |
|     | Glu Ala Th                     | _         | Glu Ser   |         |            | Leu  | Pro       | Ser   | _     | Glu   | Asp    |      |  |  |
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| 212 | ggtgcggcac                     | gaggagtti | t cccggc  | agcg ag | rgaggt     | tcct | gago      | cagca | atg ( | gadag | gagga  | 240  |  |  |
| 213 | gcgccttccc                     | tgccgccgd | g ctctgg  | ctct gg | ragcat     | tcct | cct       | tgcc  | etg ( | ctggd | cactgc | 300  |  |  |
|     | gggcggaggc                     |           |           |         |            |      |           |       |       |       |        |      |  |  |
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|     | aatgccaatg                     |           |           |         | _          |      |           | _     |       | _     |        |      |  |  |
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|     | aaataatgtt                     |           |           |         |            |      |           |       |       |       |        |      |  |  |
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|     | tagattttct                     |           |           |         |            |      |           |       |       |       |        |      |  |  |
|     | aaattttcag                     |           |           |         |            |      | _         |       |       |       |        |      |  |  |
|     | gggcagggga                     | _         |           |         |            | _    | -         | _     |       | _     |        |      |  |  |
|     | atggtgcagt                     |           |           |         |            |      |           |       |       |       |        |      |  |  |
|     | ttgttacatt                     |           |           |         |            |      |           |       |       |       |        |      |  |  |
|     | ttaccattat                     |           | _         |         |            |      |           | _     | _     | -     |        |      |  |  |
|     | ttaaacaata                     |           |           |         |            |      |           |       |       |       |        |      |  |  |
| 242 | gcattggctt                     | gaagcaata | it aatata | ttgt aa | acaaa      | aaca | cago      | tctt  | cac   | ctaat | aaaca  | 2040 |  |  |



DATE: 02/06/2002 9/904,956 TIME: 08:16:11

PATENT APPLICATION: US/09/904,956

Input Set : N:\jumbos\904956.txt

Output Set: N:\CRF3\02062002\I904956.raw

```
243 ttttatactg tttgtatgta taaaataaag gtgctgcttt agttttttgg aaaaaaaaa 2100
244 aaaaaaaaa aaaaaaaaa aaaaaaaaaa gggcggccgc gactctagag tcgacctgca 2160
245 gaagettgge egecatggee caacttgttt attgeagett ataatg
247 <210> SEQ ID NO: 4
248 <211> LENGTH: 379
249 <212> TYPE: PRT
250 <213> ORGANISM: Homo sapiens
252 <400> SEQUENCE: 4
253 Met Ala Arg Arg Ser Ala Phe Pro Ala Ala Ala Leu Trp Leu Trp Ser
                                           5
                                                                                 10
256 Ile Leu Leu Cys Leu Leu Ala Leu Arg Ala Glu Ala Gly Pro Pro Gln
                               20
                                                                         25
259 Glu Glu Ser Leu Tyr Leu Trp Ile Asp Ala His Gln Ala Arg Val Leu
                                                                40
262 Ile Gly Phe Glu Glu Asp Ile Leu Ile Val Ser Glu Gly Lys Met Ala
                                                        55
265 Pro Phe Thr His Asp Phe Arg Lys Ala Gln Gln Arg Met Pro Ala Ile
                                               70
                                                                                        75
268 Pro Val Asn Ile His Ser Met Asn Phe Thr Trp Gln Ala Ala Gly Gln
                                                                                90
                                        85
271 Ala Glu Tyr Phe Tyr Glu Phe Leu Ser Leu Arg Ser Leu Asp Lys Gly
                               100
                                                                       105
274 Ile Met Ala Asp Pro Thr Val Asn Val Pro Leu Leu Gly Thr Val Pro
                       115
                                                               120
                                                                                                       125
277 His Lys Ala Ser Val Val Gln Val Gly Phe Pro Cys Leu Gly Lys Gln
                                                      135
280 Asp Gly Val Ala Ala Phe Glu Val Asp Val Ile Val Met Asn Ser Glu
                                               150
                                                                                      155
283 Gly Asn Thr Ile Leu Gln Thr Pro Gln Asn Ala Ile Phe Phe Lys Thr
                                     165
                                                                       170 175
286 Cys Gln Gln Ala Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys
287 180
                                                                       185
289 Asn Glu Arg Arg Ile Cys Glu Cys Pro Asp Gly Phe His Gly Pro His
                                                               200
292 Cys Glu Lys Ala Leu Cys Thr Pro Arg Cys Met Asn Gly Gly Leu Cys
                                                       215
295 Val Thr Pro Gly Phe Cys Ile Cys Pro Pro Gly Phe Tyr Gly Val Asn
                                               230
                                                                                       235
298 Cys Asp Lys Ala Asn Cys Ser Thr Thr Cys Phe Asn Gly Gly Thr Cys
                                      245
                                                                               250
301 Phe Tyr Pro Gly Lys Cys Ile Cys Pro Pro Gly Leu Glu Glu Glu Gln
                               260
                                                                       265
304 Cys Glu Ile Ser Lys Cys Pro Gln Pro Cys Arg Asn Gly Gly Lys Cys
305 275
                                                               280
307 Ile Gly Lys Ser Lys Cys Lys Cys Ser Lys Gly Tyr Gln Gly Asp Leu
                                                       295
                                                                                              300
310 Cys Ser Lys Pro Val Cys Glu Pro Gly Cys Gly Ala His Gly Thr Cys
                                               310
                                                                                       315
313 His Glu Pro Asn Lys Cys Gln Cys Gln Glu Gly Trp His Gly Arg His
                                   and the first of the state of t
```

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## VERIFICATION SUMMARY

L:6540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:206

DATE: 02/06/2002

PATENT APPLICATION: US/09/904,956

TIME: 08:16:12

Input Set : N:\jumbos\904956.txt

Output Set: N:\CRF3\02062002\I904956.raw

L:34 M:270 C: Current Application Number differs, Replaced Current Application Number
L:35 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:769 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:3586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:113
L:4040 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:5344 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:174
L:5479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175